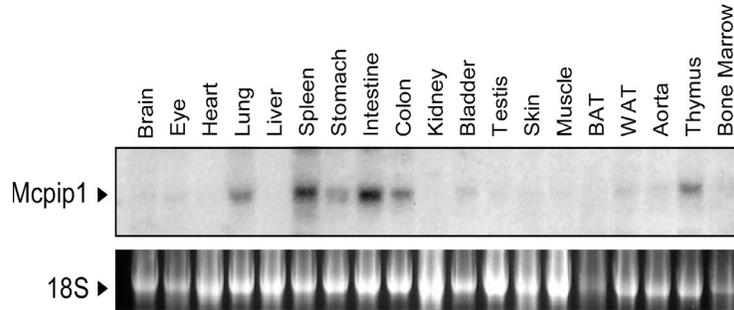
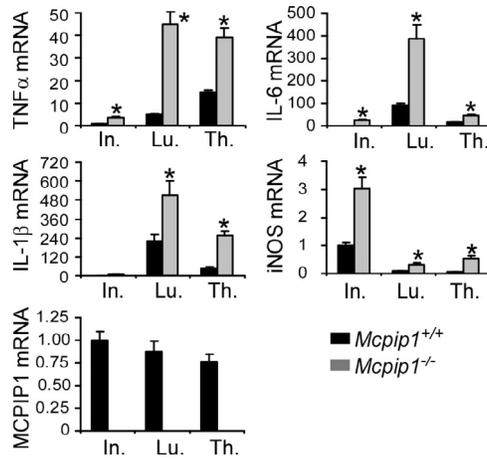


SUPPLEMENTAL MATERIAL

Liang et al., <http://www.jem.org/cgi/content/full/jem.20092641/DC1>



**Figure S1. Tissue distribution of *Mcpip1* mRNA.** A range of total RNA (10 mg/lane) extracted from adult mouse tissues as indicated was examined by RNA blot analysis. The identity of probes is shown to the left of the figure. The ethidium bromide staining image for 18S RNA served as a loading control. BAT, brown adipose tissue; WAT, white adipose tissue.



**Figure S2. Total RNA isolated from the intestine, lung, and thymus was reverse transcribed and analyzed by QPCR with the primers as indicated.** Data are presented as mean  $\pm$  SD ( $n = 3$ ; \*,  $P < 0.01$  vs. *Mcpip1*<sup>+/+</sup> group). Student's *t* test was performed. The experiment was performed on two independent sets of animals with triplicate measurements. In., intestine; Lu., lung; Th., thymus.